## SEQUENCE LISTING

- <110> Beer, Steven V. Bauer, David W.
- <120> OOMYCETE-RESISTANT TRANSGENIC PLANTS BY VIRTUE OF PATHOGEN-INDUCED EXPRESSION OF A HETEROLOGOUS HYPERSENSITIVE RESPONSE ELICITOR
- <130> 19603/2501
- <140>
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- <150> 60/178,565
- <151> 2000-01-26
- <160> 26
- <170> PatentIn Ver. 2.1
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- Gly Ala Ser Ser Lys Gly Leu Gly Met Ser Asn Gln Leu Gly Gln Ser 65 70 75 80
- Phe Gly Asn Gly Ala Gln Gly Ala Ser Asn Leu Leu Ser Val Pro Lys
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Asn Ala Phe Gly Ser Gly Val Asn Asn Ala Leu Ser Ser Ile Leu Gly 145 150 155 160

Asn Gly Leu Gly Gln Ser Met Ser Gly Phe Ser Gln Pro Ser Leu Gly 165 170 175

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Asp Lys Glu Asp Arg Gly Met Ala Lys Glu Ile Gly Gln Phe Met Asp 225 230 235 240

Gln Tyr Pro Glu Ile Phe Gly Lys Pro Glu Tyr Gln Lys Asp Gly Trp
245 250 255

Ser Ser Pro Lys Thr Asp Asp Lys Ser Trp Ala Lys Ala Leu Ser Lys 260 265 270

Pro Asp Asp Gly Met Thr Gly Ala Ser Met Asp Lys Phe Arg Gln 275 280 285

Ala Met Gly Met Ile Lys Ser Ala Val Ala Gly Asp Thr Gly Asn Thr 290 295 300

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Asn Ala

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A	sn	Ala	Gly 35	Leu	Gly	Gly	Asn	Ser 40	Ala	Leu	Gly	Leu	Gly 45	Gly	Gly	Ası
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G.	ln	Asp	Gly	Thr 180	Gln	Gly	Ser	Ser	Ser 185	Gly	Gly	Lys	Gln	Pro 190	Thr	Glı
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L	eu	Met 210	Gly	Asn	Gly	Leu	Ser 215	Gln	Leu	Leu	Gly	Asn 220	Gly	Gly	Leu	Gl
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245

Gly Gly Lys Gly Leu Gln Asn Leu Ser Gly Pro Val Asp Tyr Gln Gln

250

255

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gcctataaaa aaggagtcac tgatgcgctg tcgggcctga tgggtaatgg tctgagccag 720 ctccttggca acgggggact gggaggtggt cagggcggta atgctggcac gggtcttgac 780 ggttcgtcgc tgggcggcaa agggctgcaa aacctgagcg ggccggtgga ctaccagcag 840 ttaggtaacg ccgtgggtac cggtatcggt atgaaagcg gcattcaggc gctgaatgat 900 atcggtacgc acaggcacag ttcaacccgt tctttcgtca ataaaggcga tcgggcgatg 960 gcgaaggaaa tcggtcagta catggaccag tatcctgagg tgtttggcaa gccgcagtac 1020 cagaaaggcc cgggtcagga ggtgaaaacc gatgacaaat catgggcaaa agcactgagc 1080 aagccagatg acgacggaat ggcgaacagca agtatggagc acgcggtgcc 1200 ggtggttctt cgctgggtat tgatgccatg atggccggt atgccattaa caatatggca 1260 cttggcaagc tgggcggc ttaagct 1288

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Ser Lys Ala Leu Gln Glu Val Val Lys Leu Ala Glu Glu Leu Met
35 40 45

Arg Asn Gly Gln Leu Asp Asp Ser Ser Pro Leu Gly Lys Leu Leu Ala
50 55 60

Lys Ser Met Ala Ala Asp Gly Lys Ala Gly Gly Gly Ile Glu Asp Val 65 70 75 80

Ile Ala Ala Leu Asp Lys Leu Ile His Glu Lys Leu Gly Asp Asn Phe 85 90 95

Gly Ala Ser Ala Asp Ser Ala Ser Gly Thr Gly Gln Gln Asp Leu Met
100 105 110

Thr Gln Val Leu Asn Gly Leu Ala Lys Ser Met Leu Asp Asp Leu Leu 115 120 125

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Leu Asp Gly Asp Glu Thr Ala Ala Phe Arg Ser Ala Leu Asp Ile Ile 180 185 190

Gly Gln Gln Leu Gly Asn Gln Gln Ser Asp Ala Gly Ser Leu Ala Gly
195 200 205

Thr Gly Gly Leu Gly Thr Pro Ser Ser Phe Ser Asn Asn Ser Ser 210 215 220

Val Met Gly Asp Pro Leu Ile Asp Ala Asn Thr Gly Pro Gly Asp Ser 225 230 235 240

Gly Asn Thr Arg Gly Glu Ala Gly Gln Leu Ile Gly Glu Leu Ile Asp
245 250 255

Arg Gly Leu Gln Ser Val Leu Ala Gly Gly Gly Leu Gly Thr Pro Val 260 265 270

Asn Thr Pro Gln Thr Gly Thr Ser Ala Asn Gly Gly Gln Ser Ala Gln 275 280 285

Asp Leu Asp Gln Leu Leu Gly Gly Leu Leu Leu Lys Gly Leu Glu Ala 290 295 300

Thr Leu Lys Asp Ala Gly Gln Thr Gly Thr Asp Val Gln Ser Ser Ala 305 310 315 320

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<212> DNA

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<212> PRT

<213> Pseudomonas solanacearum

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Val Gln Asp Leu Ile Lys Gln Val Glu Lys Asp Ile Leu Asn Ile Ile 35 40 45

Ala Ala Leu Val Gln Lys Ala Ala Gln Ser Ala Gly Gly Asn Thr Gly
50 55 60

Asn Thr Gly Asn Ala Pro Ala Lys Asp Gly Asn Ala Asn Ala Gly Ala 65 70 75 80

Asn Asp Pro Ser Lys Asn Asp Pro Ser Lys Ser Gln Ala Pro Gln Ser 85 90 95

Ala Asn Lys Thr Gly Asn Val Asp Asp Ala Asn Asn Gln Asp Pro Met
100 105 110

Gln Ala Leu Met Gln Leu Leu Glu Asp Leu Val Lys Leu Leu Lys Ala 115 120 125

Ala Leu His Met Gln Gln Pro Gly Gly Asn Asp Lys Gly Asn Gly Val 130 135 140

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- Gly Ala Gly Ala Gly Gly Ala Gly Gly Val Gly Gly Ala Gly Gly
  180 185 190
- Ala Asp Gly Gly Ser Gly Ala Gly Gly Ala Gly Gly Ala Asn Gly Ala 195 200 205
- Asp Gly Gly Asn Gly Val Asn Gly Asn Gln Ala Asn Gly Pro Gln Asn 210 215 220
- Ala Gly Asp Val Asn Gly Ala Asn Gly Ala Asp Asp Gly Ser Glu Asp 225 230 235 240
- Gln Gly Gly Leu Thr Gly Val Leu Gln Lys Leu Met Lys Ile Leu Asn 245 250 255
- Ala Leu Val Gln Met Met Gln Gln Gly Gly Leu Gly Gly Gly Asn Gln 260 265 270
- Ala Gln Gly Gly Ser Lys Gly Ala Gly Asn Ala Ser Pro Ala Ser Gly 275 280 285
- Ala Asn Pro Gly Ala Asn Gln Pro Gly Ser Ala Asp Asp Gln Ser Ser 290 295 300
- Gly Gln Asn Asn Leu Gln Ser Gln Ile Met Asp Val Val Lys Glu Val 305 310 315 320
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<212> DNA

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cagtgggtga catgagcaca taagtcatct ttattgtatt ttattactca ctccaaaaat 600
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696

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